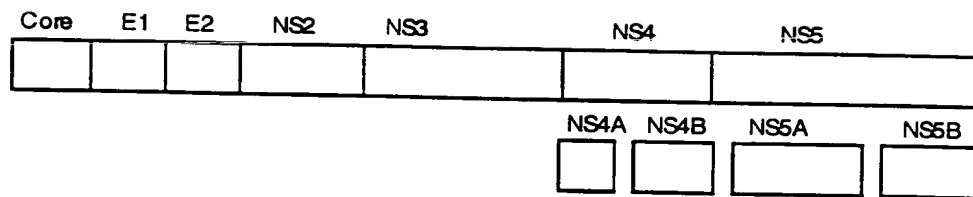


FIGURE 1



666260" 2E420460

FIGURE 2

SEQ. ID. NO:1

MSHVAVENALGLDQQFAGLDLNSSDNQSGGSTASKGRYIPPHLRNREATRGFYDKDSSGWSSSKDKDAYSSFGSRSDSRGKSSFFS  
DRGSGSRGRFDDRGRSDYDGIGSRGDRSGFGKFERGGNSRWCDKSEDDWSKPLPPSERLEQELFSGGNTGINFEKYDDIPVEATG  
NNCPPHIESFSDVEMGEIIMGNIELTRYTRPTPVQKHAIPIIKEKRDLMACAQTGSGKTA AFLLPILSQIYSDGPGEALRAMKENG  
RYGRRKQYPISLVLAPTRELAVQIYEEARKFSYRSRVRPCVVYGGADIGQQIRD LERGCHLLVATPGRLVDMMERGKIGLDFCKYL  
VLDEADRMLDMGFEPQIRRIVEQDTMPPKGV RHTMMFSATFPKEIQMLARDFLDEYIFLAVGRVGS TSENITQKVWVEESDKRSF  
LLDLLNATGKDSLTLVFVETKKGADSLEDFLYHEGYACTSIHGDRSQRDREEALHQFRSGKSPILVATAVAARGLDISNVKHVIN  
DLPSDIEEYVHRIGRTGRVGNLGLATSFNERNINITKDLLDLLEAKQEVPSWLENMAYEHYKGSSRGRSKSSRFSGGFGARDY  
ROSSGASSSSFSSSRASSSRSGGGHGSRRGFGGGYGGFYNSDGYGGNYNSQGVDDWGN

SEQ. ID. NO:2

GSTSENITQKVWVEESDKRSFLDLLNATGKDSLTLVFVETKKGADSLEDFLYHEGYACTSIHGDRSQRDREEALHQFRSGKSPI  
LVATAVAARGLDISNVKHVINFDLPSDIEEYVHRIGRTGRVGNLGLATSFNERNINITKDLLDLLEAKQEVPSWLENMAYEHY  
KGSSRGRSKSSRFSGGFGARDYRQSSGASSSSFSSSRASSSRSGGGHGSRRGFGGGYGGFYNSDGYGGNYNSQGVDDWGN

66260-2440460

FIGURE 3

SEQ. ID. NO:3

VGSTSENITQKVWVEESDKRSFLLDLLNATGKDSLTLVFVETKKGADSLEDFLYHEGYACTSIHGDRSQRDREEALHQFRSGKSP  
ILVATAVAARGLDISNVKHVINFDLPSDIEEYVHRIGRTGRVGNLGLAT

66250" 25420160

FIGURE 4

## SEQ. ID. NO:4

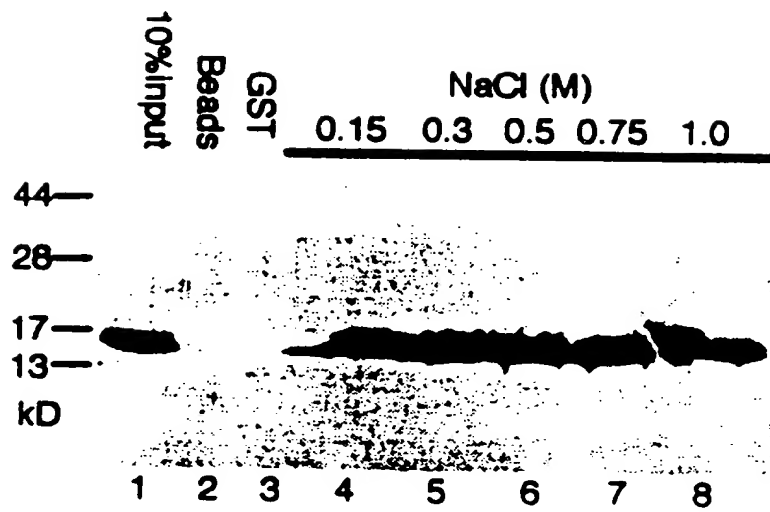
1 atggatgac gagaggatct ggtgtaccag gcgaagctgg ccgagcaggc tgagcgatac  
 61 gacgaaatgg tggagtcaat gaagaaagta gcagggatgg atgtggagct gacagttgaa  
 121 gaaagaaacc tcctatctgt tgcataataag aatgtgattg gagctagaag agcctcctgg  
 181 agaataatca gcagcattga acagaaagaa gaaaacaagg gaggagaaga caagctaaaa  
 241 atgattcggg aatatcggca aatgggttgag actgagctaa agttaatctg ttgtgacatt  
 301 ctggatgtac tggacaaaca cctcattcca gcagctaaca ctggcgagtc caagggtttc  
 361 tattataaaa tgaaagggga ctaccacagg tatctggcag aatttgccac aggaaacgac  
 421 aggaaggagg ctgcggagaa cagcctagtg gcttataaag ctgctagtga tattgcaatg  
 481 acagaacttc caccaacgca tcctattcgc ttaggtcttg ctctcaattt ttccgtattc  
 541 tactacgaaa ttcttaattc ccctgaccgt gcctgcaggt tggcaaaagc agcttttgat  
 601 gatgcaattg cagaactgga tacgtgagt gaagaaagct ataaggactc tacacttacc  
 661 atgcagttgt tacgtgataa tctgacacta tggacttcag acatgcaggg tgacggtgaa  
 721 gagcagaata aagaagcgct gcaggacgtg gaagacgaaa atcagtgaga cataagccaa  
 781 caagagaaac ca

## SEQ. ID. NO:5

translation="MDDREDLVYQAKLAEQAERYDEMVESMKKVAGMDVELTVEERNL  
 LSVAYKNVIGARRASWRIISSIEQKEENKGGEDKLKMIREYRQMVETELKLICCDILD  
 VLDKHLIPAANTGESKVFYYKMGDYHRYLAEFATGNDRKEAAENSLVAYKAASDIAM  
 TELPPTHPIRLGLALNFSVFYYEILNSPDRACRLAKAAFDDAIAELDTLSEESYKDST  
 LIMQLLRDNLTLWTSMDMQGDGEEQNKEALQDVEDENQ"

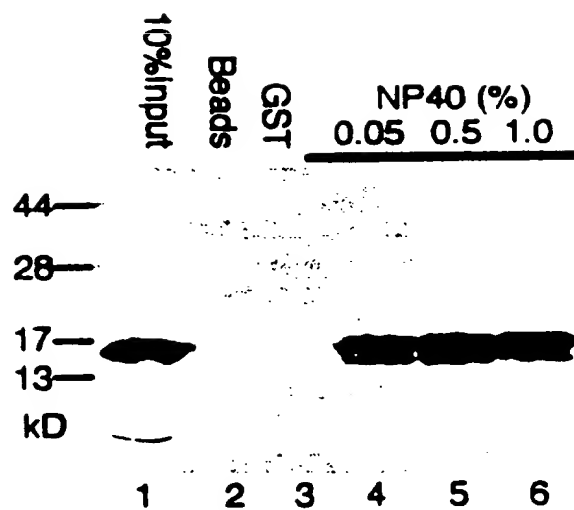
55250" 242060

FIGURE 5A



09407432.092999

FIGURE 5B



094074320450

### HCV Core Protein-DBX Interaction

[illegible]

FIGURE 6B

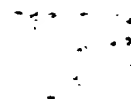
**DBX**



**PL10**



**Ded1p**



094046022666



FIGURE 7A

CORE

DBX

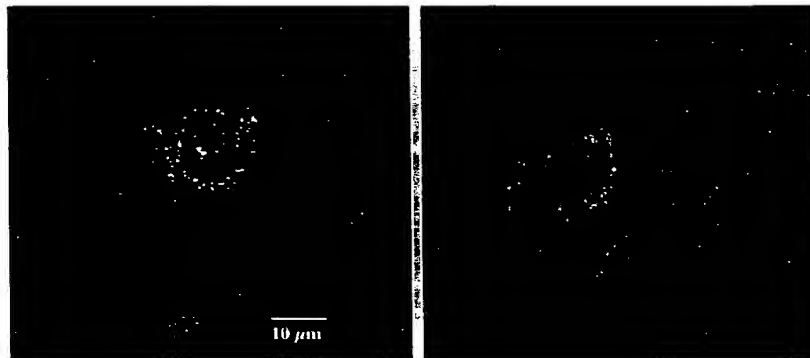
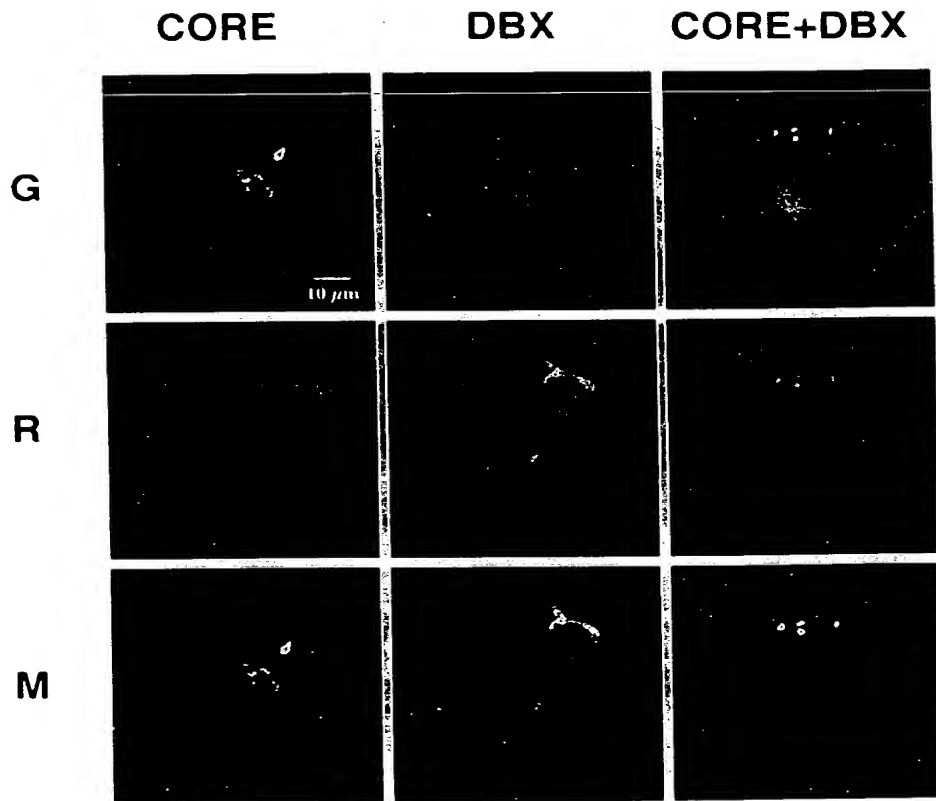
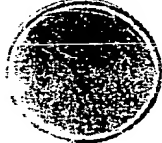
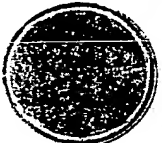
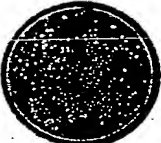
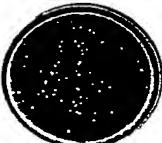
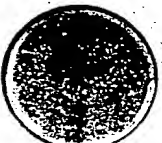



FIGURE 7B



666250 2240460

	DBX	PL10	Ded1p
CORE +			
-			

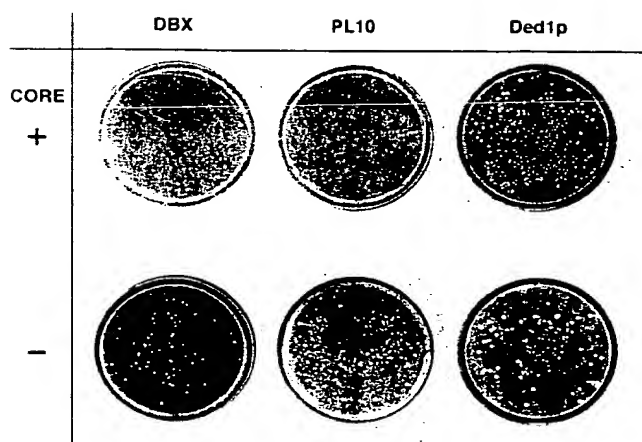


Figure 9

